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	DB=PC	SPB, USPT, EPAB, JPAB, DWPI; PLUR=YES; OP=OR	
	L6	14 and (Leonhartsberger or Maier).in.	4
	L5	L4 same synthetas\$4	48
	L4	(sam or adomet\$4) same (method\$4 or produc\$4 or synthe\$4 or biosynt\$5) same (bacter\$4 or coli\$4)	276
	L3	l2 same escheri\$4	126
	L2	L1 same synthetas\$4	488
	L1	(sam\$4 or adomet\$4) same (method\$4 or produc\$4 or synthe\$4 or biosynt\$5) same (bacter\$4 or coli\$4)	46982

END OF SEARCH HISTORY

(FILE 'HOME' ENTERED AT 20:40:17 ON 04 SEP 2006)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHOS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 20:40:48 ON 04 SEP 2006 SEA (SAM OR ADOMET?)(S)(SYNTHETAS? OR SYNTHAS?)

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48 FILE AGRICOLA
 1 FILE AQUASCI
18 FILE BIOENG
280 FILE BIOSIS
40 FILE BIOTECHABS
40 FILE BIOTECHDS
158 FILE BIOTECHNO
126 FILE CABA
354 FILE CAPLUS
5 FILE CEABA-VTB
2 FILE CONFSCI
 1 FILE CROPU
24 FILE DDFU
177 FILE DGENE
33 FILE DISSABS
39 FILE DRUGU
 5 FILE EMBAL
179 FILE EMBASE
236 FILE ESBIOBASE
6 FILE FROSTI
11 FILE FSTA
576 FILE GENBANK
32 FILE IFIPAT
22 FILE JICST-EPLUS
153 FILE LIFESCI
221 FILE MEDLINE
2 FILE NTIS
123 FILE PASCAL
2 FILE PROMT
215 FILE SCISEARCH
132 FILE TOXCENTER
299 FILE USPATFULL
23 FILE USPAT2
24 FILE WPIDS
24 FILE WPINDEX
2 FILE NLDB
  QUE (SAM OR ADOMET?)(S)(SYNTHETAS? OR SYNTHAS?)
 D RANK
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FILE 'CAPLUS, USPATFULL, BIOSIS, ESBIOBASE, MEDLINE, SCISEARCH, EMBASE, BIOTECHNO, LIFESCI, TOXCENTER' ENTERED AT 20:42:47 ON 04 SEP 2006 2227 SEA (SAM OR ADOMET?)(S)(SYNTHETAS? OR SYNTHAS?)

- L2
- L3 1715 SEA L2(S)(METHOD? OR SYNTHE? OR BIOSYNTHE? OR PRODUC? OR PROCES?)
- 359 SEA L3(S)(BACTER? OR COLI? OR ESCHERI?) L4
- L5 176 DUP REM L4 (183 DUPLICATES REMOVED)

D TI L5 1-175

L1

D IBIB ABS L5 39 40 54 111 114-115 143 146 169

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=> index bioscience medicine FILE 'DRUGMONOG' ACCESS NOT AUTHORIZED COST IN U.S. DOLLARS

SINCE FILE TOTAL ENTRY SESSION 0.21 0.21

FULL ESTIMATED COST

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 20:40:48 ON 04 SEP 2006

71 FILES IN THE FILE LIST IN STNINDEX

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```
=> s (sam or adomet?)(s)(synthetas? or synthas?)
              FILE AGRICOLA
         48
              FILE AQUASCI
          1
              FILE BIOENG
         18
        280
              FILE BIOSIS
         40
              FILE BIOTECHABS
         40
              FILE BIOTECHDS
              FILE BIOTECHNO
        158
              FILE CABA
        126
              FILE CAPLUS
        354
          5
              FILE CEABA-VTB
          2
              FILE CONFSCI
              FILE CROPU
          1
         24
              FILE DDFU
              FILE DGENE
        177
         33
              FILE DISSABS
         39
              FILE DRUGU
  27 FILES SEARCHED...
          5
              FILE EMBAL
        179
              FILE EMBASE
        236
              FILE ESBIOBASE
          6
              FILE FROSTI
         11
              FILE FSTA
        576
              FILE GENBANK
         32
              FILE IFIPAT
         22
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              FILE LIFESCI
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              FILE MEDLINE
              FILE NTIS
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        123
              FILE PASCAL
          2
              FILE PROMT
        215
              FILE SCISEARCH
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              FILE TOXCENTER
        299
              FILE USPATFULL
         23
              FILE USPAT2
  64 FILES SEARCHED...
         24
              FILE WPIDS
         24
              FILE WPINDEX
              FILE NLDB
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- 36 FILES HAVE ONE OR MORE ANSWERS, 71 FILES SEARCHED IN STNINDEX
- L1 QUE (SAM OR ADOMET?) (S) (SYNTHETAS? OR SYNTHAS?)

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=> d rank
F1
          576
                GENBANK
F2
          354
                CAPLUS
F3
          299
                USPATFULL
F4
          280
                BIOSIS
F5
          236
                ESBIOBASE
F6
          221
                MEDLINE
F7
          215
                SCISEARCH
F8
          179
                EMBASE
F9
          177 DGENE
          158 BIOTECHNO
F10
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F11	153	LIFESCI
F12	132	TOXCENTER
F13	126	CABA
F14	123	PASCAL
F15	48	AGRICOLA
F16	40	BIOTECHABS
F17	40	BIOTECHDS
F18	39	DRUGU
F19	33	DISSABS
F20	32	IFIPAT
F21	24	DDFU
F22	24	WPIDS
F23	24	WPINDEX
F24	23	USPAT2
F25	22	JICST-EPLUS
F26	18	BIOENG
F27	11	FSTA
F28	6	FROSTI
F29	5	CEABA-VTB
F30	5	EMBAL
F31	2	CONFSCI
F32	2	NTIS
F33	2	PROMT
F34	2	NLDB
F35	1	AQUASCI
F36	1	CROPU

=> file f2-f8, f10-12 COST IN U.S. DOLLARS

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FULL ESTIMATED COST

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=> s (sam or adomet?)(s)(synthetas? or synthas?)
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- L2 2227 (SAM OR ADOMET?) (S) (SYNTHETAS? OR SYNTHAS?)
- => s 12(s) (method? or synthe? or biosynthe? or produc? or proces?)
 - 4 FILES SEARCHED...
- L3 1715 L2(S) (METHOD? OR SYNTHE? OR BIOSYNTHE? OR PRODUC? OR PROCES?)
- => s l3(s)(bacter? or coli? or escheri?)
- 8 FILES SEARCHED...
- L4 359 L3(S) (BACTER? OR COLI? OR ESCHERI?)
- => dup rem 14
- PROCESSING COMPLETED FOR L4
- L5 176 DUP REM L4 (183 DUPLICATES REMOVED)
- => d ti 15 1-175
- L5 ANSWER 1 OF 176 USPATFULL on STN
- TI Methods and systems for predicting cancer outcome
- L5 ANSWER 2 OF 176 USPATFULL on STN
- TI Methods for predicting cancer outcome and gene signatures for use therein
- L5 ANSWER 3 OF 176 USPATFULL on STN
- TI Methods for indentifying compounds that modulate an enzyme involved in riboflavin metabolism in a pathogenic microorganism
- L5 ANSWER 4 OF 176 USPATFULL on STN
- TI Methods for indentifying compounds that modulate an enzyme involved in reductive carboxylation in a pathogenic microorganism
- L5 ANSWER 5 OF 176 USPATFULL on STN
- TI Methods for indentifying compounds that modulate an enzyme involved in thiamine metabolism in a pathogenic microorganism
- L5 ANSWER 6 OF 176 USPATFULL on STN
- TI Pharmaceutical compositions
- L5 ANSWER 7 OF 176 USPATFULL on STN
- TI Inhibitors of nucleoside phosphorylases and nucleosidases
- L5 ANSWER 8 OF 176 USPATFULL on STN
- TI Feedback-resistant homoserine transsuccinylases having a modified c terminus
- L5 ANSWER 9 OF 176 USPATFULL on STN
- TI Methods for indentifying compounds that modulate an enzyme in the coenzyme a biosynthetic pathway in a pathogenic microoganism
- L5 ANSWER 10 OF 176 USPATFULL on STN
- TI Nanomachine compositions and methods of use
- L5 ANSWER 11 OF 176 USPATFULL on STN
- TI Methods of identifying patients at risk of developing encephalitis following immunotherapy for Alzheimer's disease
- L5 ANSWER 12 OF 176 USPATFULL on STN
- TI Complete genome and protein sequence of the hyperthermophile methanopyrus kandleri av19 and monophyly of archael methanogens and methods of use thereof
- L5 ANSWER 13 OF 176 USPATFULL on STN
- TI Method for the identification and treatment of pathogenic microorganism infections by inhibiting one or more enzymes in an essential metabolic

pathway

- L5 ANSWER 14 OF 176 USPATFULL on STN
- TI Novel full length cDNA
- L5 ANSWER 15 OF 176 USPATFULL on STN
- TI Nucleic acid sequences relating to Bacteroides fragilis for diagnostics and therapeutics
- L5 ANSWER 16 OF 176 USPATFULL on STN
- Nucleic acid and amino acid sequences relating to Staphylococcus epidermidis for diagnostics and therapeutics
- L5 ANSWER 17 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Identification of fungal sphingolipid C9-methyltransferases by phylogenetic profiling
- L5 ANSWER 18 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI In vivo hydrolysis of S-adenosylmethionine induces the met regulon of Escherichia coli
- L5 ANSWER 19 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI In vivo hydrolysis of S-adenosylmethionine induces the met regulon of Escherichia coli
- L5 ANSWER 20 OF 176 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
- TI Assessment of inhibitors of S-adenosylmethionine synthesis discovered by computational docking.
- L5 ANSWER 21 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI The Structure of the RNA m.sup.5C Methyltransferase YebU from Escherichia coli Reveals a C-terminal RNA-recruiting PUA Domain
- L5 ANSWER 22 OF 176 MEDLINE on STN
- TI Molecular evolution of AdoMet synthetase by DNA recombination with a novel separate-mixing method.
- L5 ANSWER 23 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 2
- TI Molecular evolution of adomet synthetase by DNA recombination with a novel Separate-Mixing method
- L5 ANSWER 24 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Characterization of S-adenosylmethionine synthetase from Streptomyces avermitilis NRRL8165 and its effect on antibiotic production
- L5 ANSWER 25 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 4
- TI Cloning and expression of rat liver S-adenosylmethionine synthetase
- L5 ANSWER 26 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 5
- TI The SMK box is a new SAM-binding RNA for translational regulation of SAM synthetase
- L5 ANSWER 27 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
- TI Spectral and kinetic characterization of 7,8-diaminopelargonic acid synthase from Mycobacterium tuberculosis
- L5 ANSWER 28 OF 176 USPATFULL on STN
- TI Constructs and methods for the regulation of gene expression

- L5 ANSWER 29 OF 176 USPATFULL on STN
- TI Corynebacterium glutamicum genes encoding proteins involved in membrane synthesis and membrane transport
- L5 ANSWER 30 OF 176 USPATFULL on STN
- TI Quorum sensing signaling in bacteria
- L5 ANSWER 31 OF 176 USPATFULL on STN
- TI Nucleic acid and amino acid sequences relating to streptococcus pneumoniae for diagnostics and therapeutics
- L5 ANSWER 32 OF 176 USPATFULL on STN
- TI Adenosylmethionine synthetase from streptomyces sp gene sequences coding the same and method for mass production for secondary metabolites including antibiotics thereof
- L5 ANSWER 33 OF 176 USPATFULL on STN
- TI Identification and isolation of novel polypeptides having PDZ domains and methods of using same
- L5 ANSWER 34 OF 176 USPATFULL on STN
- TI Protein interaction mapping
- L5 ANSWER 35 OF 176 USPATFULL on STN
- TI Quorum sensing signaling in bacteria
- L5 ANSWER 36 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI A nucleosidase required for in vivo function of the S-adenosyl-L-methionine radical enzyme, biotin synthase
- L5 ANSWER 37 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Transcriptional analysis of the cyclopropane fatty acid synthase gene of Lactococcus lactis MG1363 at low pH
- L5 ANSWER 38 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Developmental regulation of 1-aminocyclopropane-1-carboxylate synthase gene expression during leaf ontogeny in white clover
- L5 ANSWER 39 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 8
- TI Method for fermentative preparation of S-adenosylmethionine using recombinant Escherichia coli
- L5 ANSWER 40 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN
- TI Enzymatic catalysis method for producing S-adenosylmethionine
- L5 ANSWER 41 OF 176 USPATFULL on STN
- TI Inhibitors of spermidine synthase for the treatment of osteoarthritis and cartilage rehabilitation
- L5 ANSWER 42 OF 176 USPATFULL on STN
- TI Polypeptides involved in the biosynthesis of streptogramins, nucleotide sequences coding for these polypeptides and their use
- L5 ANSWER 43 OF 176 USPATFULL on STN
- TI Nucleic acid molecules encoding proteins essential for plant growth and development and uses thereof
- L5 ANSWER 44 OF 176 USPATFULL on STN
- TI Treatment of patients with multiple sclerosis based on gene expression changes in central nervous system tissues
- L5 ANSWER 45 OF 176 USPATFULL on STN

- TI Overcoming DAPA aminotransferase bottlenecks in biotin vitamers biosynthesis
- L5 ANSWER 46 OF 176 USPATFULL on STN
- TI Screening method for anti-microbial drug targets by genome-saturating mutagenesis (gsm)
- L5 ANSWER 47 OF 176 USPATFULL on STN
- TI Methods and compositions for determining enzymatic activity and specificity of methlytransferases
- L5 ANSWER 48 OF 176 USPATFULL on STN
- TI Fungal gene cluster associated with pathogenesis
- L5 ANSWER 49 OF 176 USPATFULL on STN
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- TI Senescence-associated plant promotors
- L5 ANSWER 51 OF 176 USPATFULL on STN
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- L5 ANSWER 52 OF 176 USPATFULL on STN
- TI Quorum sensing signaling in bacteria
- L5 ANSWER 53 OF 176 USPATFULL on STN
- TI Human cystathionine beta-synthase variants and methods of production thereof
- L5 ANSWER 54 OF 176 USPATFULL on STN
- TI Corynebacterium glutamicum genes encoding proteins involved in membrane synthesis and membrane transport
- L5 ANSWER 55 OF 176 USPATFULL on STN
- TI Streptococcus pneumoniae polynucleotides and sequences
- L5 ANSWER 56 OF 176 USPATFULL on STN
- TI Novel human polynucleotides and polypeptides encoded thereby
- L5 ANSWER 57 OF 176 USPATFULL on STN
- Method for the identification and treatment of pathogenic microorganism infections by inhibiting one or more enzymes in an essential metabolic pathway and compounds and pharmaceutical compositions useful therefor
- L5 ANSWER 58 OF 176 USPATFULL on STN
- TI Methods of diagnosis of ovarian cancer, compositions and methods of screening for modulators of ovarian cancer
- L5 ANSWER 59 OF 176 USPATFULL on STN
- TI Novel full-length cDNA
- L5 ANSWER 60 OF 176 USPATFULL on STN
- TI Polypeptides from Chlamydia pneumoniae and their use in the diagnosis, prevention and treatment of disease
- L5 ANSWER 61 OF 176 USPATFULL on STN
- TI Nucleic acid and amino acid sequences relating to Streptococcus pneumoniae for diagnostics and therapeutics
- L5 ANSWER 62 OF 176 USPATFULL on STN
- TI Cathepsin V-like polypeptides
- L5 ANSWER 63 OF 176 USPATFULL on STN
- TI Nucleic acid sequences relating to Candida albicans for diagnostics and

therapeutics

- L5 ANSWER 64 OF 176 USPATFULL on STN
- TI Nucleic acids and polypeptides
- L5 ANSWER 65 OF 176 USPATFULL on STN
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- TI Tumor necrosis factor receptor 2
- L5 ANSWER 68 OF 176 USPATFULL on STN
- TI Genes expressed in C3A liver cell cultures treated with steroids
- L5 ANSWER 69 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI Escherichia coli Lipoyl Synthase Binds Two Distinct [4Fe-4S] Clusters per Polypeptide
- L5 ANSWER 70 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI Escherichia coli cyclopropane fatty acid synthase: Mechanistic and site-directed mutagenetic studies
- L5 ANSWER 71 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Role of the [2Fe-2S] Cluster in Recombinant Escherichia coli Biotin Synthase
- L5 ANSWER 72 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Characterization of the Cofactor Composition of Escherichia coli Biotin Synthase
- L5 ANSWER 73 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V.
- TI Crystal Structure of the S-Adenosylmethionine Synthetase Ternary Complex:
 A Novel Catalytic Mechanism of S-Adenosylmethionine Synthesis from ATP
 and Met
- L5 ANSWER 74 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI 5'-Methylthioadenosine Modulates the Inflammatory Response to Endotoxin in Mice and in Rat Hepatocytes
- L5 ANSWER 75 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 11
- TI Gene switching in Amoeba proteus caused by endosymbiotic bacteria
- L5 ANSWER 76 OF 176 USPATFULL on STN
- TI Compositions and methods for identifying and distinguishing orthosomycin biosynthetic loci
- L5 ANSWER 77 OF 176 USPATFULL on STN
- TI Novel nucleic acids and polypeptides
- L5 ANSWER 78 OF 176 USPATFULL on STN
- TI Differentially-regulated prostate cancer genes
- L5 ANSWER 79 OF 176 USPATFULL on STN
- TI Method for the identification and treatment of pathogenic microorganism infections by inhibiting one or more enzymes in an essential metabolic pathway

- L5 ANSWER 80 OF 176 USPATFULL on STN
- TI Genes and proteins for the biosynthesis of polyketides
- L5 ANSWER 81 OF 176 USPATFULL on STN
- TI Inhibitors of autoinducer transporters
- L5 ANSWER 82 OF 176 USPATFULL on STN
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- L5 ANSWER 83 OF 176 USPATFULL on STN
- TI Nanomachine compositions and methods of use
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- TI Structural basis of quorum sensing signal generation and methods and therapeutic agents derived therefrom
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- TI Expressed sequences of arabidopsis thaliana
- L5 ANSWER 86 OF 176 USPATFULL on STN
- TI Betaines as adjuvants to susceptibility testing and antimicrobial therapy
- L5 ANSWER 87 OF 176 USPATFULL on STN
- TI Identification of modulatory molecules using inducible promoters
- L5 ANSWER 88 OF 176 USPATFULL on STN
- TI Genes and proteins for the biosynthesis of anthramycin
- L5 ANSWER 89 OF 176 USPATFULL on STN
- TI SAM operon
- L5 ANSWER 90 OF 176 USPATFULL on STN
- TI Polypeptides involved in the biosynthesis of streptogramins, nucleotide sequences coding for these polypeptides and their use
- L5 ANSWER 91 OF 176 USPATFULL on STN
- TI Nucleic acid and amino acid sequences relating to Enterococcus faecalis for diagnostics and therapeutics
- L5 ANSWER 92 OF 176 USPATFULL on STN
- TI Nucleic acid and amino acid sequences relating to Acinetobacter baumannii for diagnostics and therapeutics
- L5 ANSWER 93 OF 176 USPATFULL on STN
- TI Nucleic acid and amino acid sequences relating to pseudomonas aeruginosa for diagnostics and therapeutics
- L5 ANSWER 94 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI Biochemical Diversity among the 1-Amino-cyclopropane-1-Carboxylate Synthase Isozymes Encoded by the Arabidopsis Gene Family
- L5 ANSWER 95 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Identification and functional reconstitution of yeast mitochondrial carrier for S-adenosylmethionine
- L5 ANSWER 96 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Transcription termination control of the S box system: Direct measurement of S-adenosylmethionine by the leader RNA
- L5 ANSWER 97 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN

- TI Control of adenosylmethionine-dependent radical generation in biotin synthase: A kinetic and thermodynamic analysis of substrate binding to active and inactive forms of BioB
- L5 ANSWER 98 OF 176 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Accumulation of S-adenosyl-L-methionine enhances production of actinorhodin but inhibits sporulation in Streptomyces lividans TK23
- L5 ANSWER 99 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN
- TI Conformational dynamics of the active site loop of S-adenosylmethionine synthetase illuminated by site-directed spin labeling
- ANSWER 100 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Bacterial-like energy metabolism in the amitochondriate protozoon Hexamita inflata
- L5 ANSWER 101 OF 176 USPATFULL on STN
- TI Inhibitors of spermidine synthase for the treatment of osteoarthritis and cartilage rehabilitation
- L5 ANSWER 102 OF 176 USPATFULL on STN
- TI Melon promoters for expression of transgenes in plants
- L5 ANSWER 103 OF 176 USPATFULL on STN
- TI OVERCOMING DAPA AMINOTRANSFERASE BOTTLENECKS IN BIOTIN VITAMERS BIOSYNTHESIS
- L5 ANSWER 104 OF 176 USPATFULL on STN
- TI Expressed sequences of arabidopsis thaliana
- L5 ANSWER 105 OF 176 USPATFULL on STN
- TI Betaines as adjuvants to susceptibility testing and antimicrobial therapy
- L5 ANSWER 106 OF 176 USPATFULL on STN
- TI Apple promoters for expression of transgenes in plants
- L5 ANSWER 107 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Enzymatic properties of S-adenosylmethionine synthetase from the archaeon Methanococcus jannaschii
- L5 ANSWER 108 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI The active site loop of S-adenosylmethionine synthetase modulates catalytic efficiency
- L5 ANSWER 109 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Studies on the role of the metK gene product of Escherichia coli K-12
- L5 ANSWER 110 OF 176 USPATFULL on STN
- TI Cotton modification using ovary-tissue transcriptional factors
- L5 ANSWER 111 OF 176 USPATFULL on STN
- TI SAM operon
- L5 ANSWER 112 OF 176 USPATFULL on STN
- Polypeptides involved in the biosynthesis of streptogramins, nucleotide sequences coding for these polypeptides and their use
- L5 ANSWER 113 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE

- TI In vitro reconstitution of the Pseudomonas aeruginosa nonribosomal peptide synthesis of pyochelin: Characterization of backbone tailoring thiazoline reductase and N-methyltransferase activities
- L5 ANSWER 114 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 19
- TI Lowering S-adenosylmethionine levels in Escherichia coli modulates C-to-T transition mutations
- L5 ANSWER 115 OF 176 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN
- TI metK is an essential gene in E. coli K-12.
- L5 ANSWER 116 OF 176 USPATFULL on STN
- TI Polypeptides involved in the biosynthesis of streptogramins, nucleotide sequences coding for these polypeptides and their use
- L5 ANSWER 117 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI Iron-sulfur cluster interconversions in biotin synthase: Dissociation and reassociation of iron during conversion of [2Fe-2S] to [4Fe-4S] clusters
- L5 ANSWER 118 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 20
- TI Energetics of S-Adenosylmethionine Synthetase Catalysis
- L5 ANSWER 119 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 21
- TI The bifunctional active site of S-adenosylmethionine synthetase. Roles of the basic residues
- L5 ANSWER 120 OF 176 USPATFULL on STN
- TI Polypeptides involved in the biosynthesis of streptogramins, nucleotide sequences coding for these polypeptides and their use
- L5 ANSWER 121 OF 176 USPATFULL on STN
- TI Combinations and methods for reducing antimicrobial resistance
- L5 ANSWER 122 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 22
- TI The bifunctional active site of S-adenosylmethionine synthetase. Roles of the active site aspartates
- L5 ANSWER 123 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 23
- TI Influence of S-adenosylmethionine pool size on spontaneous mutation, Dam methylation, and cell growth of Escherichia coli
- L5 ANSWER 124 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 24
- TI The Conformations of a Substrate and a Product Bound to the Active Site of S-Adenosylmethionine Synthetase
- L5 ANSWER 125 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 25
- TI Genome degradation is an ongoing process in Rickettsia evolution
- L5 ANSWER 126 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Molecular characterization of Plasmodium falciparum S-adenosylmethionine synthetase
- ANSWER 127 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Characterization of recombinant Arabidopsis thaliana threonine synthase
- L5 ANSWER 128 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI The biosynthesis of mycolic acids in Mycobacterium tuberculosis: Enzymatic methyl(ene) transfer to acyl carrier protein bound meromycolic acid in vitro

- L5 ANSWER 129 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 29
- TI The Active-Site Arginine of S-Adenosylmethionine Synthetase Orients the Reaction Intermediate
- L5 ANSWER 130 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI The mechanism of adenosylmethionine-dependent activation of methionine synthase: A rapid kinetic analysis of intermediates in reductive methylation of cob(II)alamin enzyme
- L5 ANSWER 131 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 30
- TI Lack of S-adenosylmethionine results in a cell division defect in Escherichia coli
- L5 ANSWER 132 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 31
- TI A new series of cyclic amino acids as inhibitors of S-adenosyl L-methionine synthetase
- L5 ANSWER 133 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI A new series of S-adenosyl-L-methionine synthetase inhibitors
- L5 ANSWER 134 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI Trypsin cleavage of human cystathionine β -synthase into an evolutionarily conserved active core: Structural and functional consequences
- L5 ANSWER 135 OF 176 USPATFULL on STN
- TI Screening for mutations by expressing cDNA segments
- L5 ANSWER 136 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN
- TI Evidence for symbiont-induced alteration of a host's gene expression: irreversible loss of SAM synthetase from Amoeba proteus
- L5 ANSWER 137 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 32
- TI Regulation of rat liver S-adenosylmethionine synthetase during septic shock: role of nitric oxide
- L5 ANSWER 138 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Nucleotide sequence and developmental expression of Acanthamoeba S-adenosylmethionine synthetase gene
- L5 ANSWER 139 OF 176 USPATFULL on STN
- TI Genetic control of ethylene biosynthesis in plants
- L5 ANSWER 140 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI A novel bifunctional fusion enzyme catalyzing ethylene synthesis via 1-aminocyclopropane-1-carboxylic acid
- L5 ANSWER 141 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Generation of cell-to-cell signals in quorum sensing: Acyl homoserine lactone synthase activity of a purified Vibrio fischeri LuxI protein
- L5 ANSWER 142 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Structure and function of S-adenosylmethionine synthetase: Crystal structures of S-adenosylmethionine synthase with ADP, BrADP, and PP(i) at 2.8 Å resolution
- L5 ANSWER 143 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 37
- TI Enzymic synthesis of S-adenosyl-L-methionine on the preparative scale

- L5 ANSWER 144 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI The structure of the C-terminal domain of methionine synthase: Presenting S-adenosylmethionine for reductive methylation of B.sub.1.sub.2
- L5 ANSWER 145 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI Flexible loop in the structure of S-adenosylmethionine synthetase crystallized in the tetragonal modification
- L5 ANSWER 146 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN
- TI Production of s-adenosyl-methionine (SAM) by fermentation of Escherichia coli transformed with rat S-Adenosyl-methionine synthetase
- L5 ANSWER 147 OF 176 USPATFULL on STN
- TI Genetic control of ethylene biosynthesis in plants using S-adenosylmethionine hydrolase
- L5 ANSWER 148 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 38
- TI Structural and functional roles of cysteine 90 and cysteine 240 in S-adenosylmethionine synthetase
- L5 ANSWER 149 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 39
- TI Investigation of monovalent cation activation of S-adenosylmethionine synthetase using mutagenesis and uranyl inhibition
- L5 ANSWER 150 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI Three cDNAs encoding S-adenosyl-L-methionine synthetase from Actinidia chinensis
- L5 ANSWER 151 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Hyperhomocysteinemia in premature arterial disease: Examination of cystathionine β -synthase alleles at the molecular level
- L5 ANSWER 152 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI Structural and functional roles of cysteine 90 and cysteine 240 in S-adenosylmethionine synthetase
- L5 ANSWER 153 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI Investigation of monovalent cation activation of S-adenosylmethionine synthetase using mutagenesis and uranyl inhibition
- L5 ANSWER 154 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Expression of apple 1-aminocyclopropane-1-carboxylate synthase in Escherichia coli: Kinetic characterization of wild-type and active-site mutant forms
- L5 ANSWER 155 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN
- TI Nitrous oxide inactivation of cobalamin-dependent methionine synthase from Escherichia coli: Characterization of the damage to the enzyme and prosthetic group
- L5 ANSWER 156 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 42
- TI Transfer and isomerization of the ribose moiety of AdoMet during the biosynthesis of queuosine tRNAs, a new unique reaction catalyzed by the QueA protein from Escherichia coli
- L5 ANSWER 157 OF 176 Elsevier BIOBASE COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Bacterial expression of catalytically active fragments of the

multifunctional enzyme enniatin synthetase

- L5 ANSWER 158 OF 176 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Isozymes of S-adenosylmethionine synthetase are encoded by tandemly duplicated genes in Escherichia coli
- L5 ANSWER 159 OF 176 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- Isolation and characterization of Escherichia coli mutants affected in aerobic respiration: The cloning and nucleotide sequence of ubiG.

 Identification of an S-adenosylmethionine-binding motif in protein, RNA, and small-molecule methyltransferases
- L5 ANSWER 160 OF 176 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN A functional tomato ACC synthase expressed in Escherichia coli demonstrates suicidal inactivation by its substrate S-adenosylmethionine
- L5 ANSWER 161 OF 176 USPATFULL on STN
- TI Lipophilic salts of S-adenosyl-L-methionine (SAM) with acylated taurine derivatives
- L5 ANSWER 162 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN
- TI Demonstration of extrinsic DNA from immune complexes in plasma of a patient with systemic lupus erythematosus
- L5 ANSWER 163 OF 176 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Novel Escherichia coli K-12 mutants impaired in S-adenosylmethionine synthesis
- L5 ANSWER 164 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN
- TI Specificity of S-adenosylmethionine synthetase for ATP analogs mono- and disubstituted in bridging positions of the polyphosphate chain
- L5 ANSWER 165 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 47
- TI Antigenic conservation of primary structural regions of S-adenosylmethionine synthetase
- L5 ANSWER 166 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN
- TI Cloning the mRNA encoding 1-aminocyclopropane-1-carboxylate synthase, the key enzyme for ethylene biosynthesis in plants.
- L5 ANSWER 167 OF 176 USPATFULL on STN
- TI Stable salts of S-adenosyl-L-methionine with polyanions
- L5 ANSWER 168 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 48
- TI SAM1, the structural gene for one of the S-adenosylmethionine synthetases in Saccharomyces cerevisiae. Sequence and expression
- L5 ANSWER 169 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 49
- TI A mutant of Escherichia coli temperature sensitive in the biosynthesis of S-adenosylmethionine
- L5 ANSWER 170 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 50
- TI S-Adenosylmethionine: studies on chemical and enzymic synthesis
- L5 ANSWER 171 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 51
- TI Characterization of the monovalent cation activator binding site of S-adenosylmethionine synthetase by thallium-205 NMR of enzyme-bound thallium(1+)
- L5 ANSWER 172 OF 176 BIOTECHNO COPYRIGHT 2006 Elsevier Science B.V. on STN DUPLICATE
- TI Regulation of methionine synthesis in Escherichia coli: Effect of metJ

gene product and S-adenosylmethionine on the in vitro expression of the metB, metL and metJ genes

L5 ANSWER 173 OF 176 LIFESCI COPYRIGHT 2006 CSA on STN

TI Structure of the divalent metal ion activator binding site of S-adenosylmethionine synthetase studied by vanadyl(IV) electron paramagnetic resonance.

L5 ANSWER 174 OF 176 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on STN DUPLICATE 53

TI S ADENOSYL METHIONINE SYNTHETASE FROM ESCHERICHIA-COLI.

L5 ANSWER 175 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 54

TI De novo synthesis of methionine in normal and brugia-infected Aedes aegypti

=> d ibib abs 15 39 40 54 111 114-115 143 146 169

L5 ANSWER 39 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 8

ACCESSION NUMBER:

2004:739861 CAPLUS

DOCUMENT NUMBER:

141:255490

TITLE:

Method for fermentative preparation of

S-adenosylmethionine using recombinant Escherichia

coli

INVENTOR(S):

Leonhartsberger, Susanne; Maier, Thomas

PATENT ASSIGNEE(S):

Consortium fur Electrochemische Industrie G.m.b.H.,

Germany

SOURCE:

U.S. Pat. Appl. Publ., 14 pp.

CODEN: USXXCO

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 2004175805	A1	20040909	US 2004-789493	20040227
DE 10309856	A1	20040923	DE 2003-10309856	20030306
CA 2457423	AA	20040906	CA 2004-2457423	20040303
EP 1457569	A1	20040915	EP 2004-5193	20040304
R: AT, BE, CH,	DE, DK	, ES, FR, GB,	, GR, IT, LI, LU, NL, SI	E, MC, PT,
IE, SI, LT,	LV, FI	, RO, MK, CY,	, AL, TR, BG, CZ, EE, HT	J, PL, SK
JP 2004267209	A2	20040930	JP 2004-62766	20040305
CN 1570126	A	20050126	CN 2004-10007493	20040305
PRIORITY APPLN. INFO.:			DE 2003-10309856 A	20030306
AB A method for ferment	tative p	production of	S-adenosylmethionine	(
SAM), includes cult	uring a	bacterial st	rain obtainable	
from a starting stra	ain and	having incre	eased SAM-	
			carting strain, in a cul	lture
medium, the bacteria				
culture medium and				
			methionine adenosyltran	nsferase, was

L5 ANSWER 40 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN

overexpressed in Escherichia coli.

ACCESSION NUMBER:

2005:66408 CAPLUS

DOCUMENT NUMBER:

142:428865

TITLE:

Enzymatic catalysis method for producing

S-adenosylmethionine

INVENTOR(S):

Zhang, Keqin

PATENT ASSIGNEE(S):

Yinsai Biological Science and Technology Co., Ltd.,

Guangzhou, Peop. Rep. China

SOURCE:

Faming Zhuanli Shenqing Gongkai Shuomingshu, 9 pp.

CODEN: CNXXEV

DOCUMENT TYPE:

Patent Chinese

LANGUAGE:

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
		~		
CN 1483829	Α	20040324	CN 2003-126834	20030612
PRIORITY APPLN. INFO.:			CN 2003-126834	20030612

The method comprises constructing the S-adenosylmethionine AΒ synthetase-gene carrying genetically engineered bacterial

strain via DNA recombination technique and enzyme engineering technique,

fermenting to obtain SAM synthetase solution;

synthesizing SAM from DL-methionine and ATP in 0.05-1 M phosphate buffer (pH 8.0) in the presence of the SAM

synthetase solution catalyst at 30-35°C, separating and purifying

on cationic exchange resin (DK110 or HD-2) column, adding hydroxypropyl Me cellulose in eluent, and spray drying.

ANSWER 54 OF 176 USPATFULL on STN

ACCESSION NUMBER:

2004:39574 USPATFULL

TITLE:

Corynebacterium glutamicum genes encoding proteins involved in membrane synthesis and membrane transport

INVENTOR(S):

Pompejus, Markus, Waldsee, GERMANY, FEDERAL REPUBLIC OF

Kroger, Burkhard, Limburgerhof, GERMANY, FEDERAL

REPUBLIC OF

Schroder, Hartwig, Nussloch, GERMANY, FEDERAL REPUBLIC

Zelder, Oskar, Speyer, GERMANY, FEDERAL REPUBLIC OF Haberhauer, Gregor, Limburgerhof, GERMANY, FEDERAL

REPUBLIC OF

PATENT ASSIGNEE(S):

BASF Aktiengesellschaft, Ludwigshafen, GERMANY, FEDERAL

REPUBLIC OF (non-U.S. corporation)

NUMBER	KIND	DATE
US 2004030116	71	20040212
US 2004030116	Al Al	20040212

PATENT INFORMATION: APPLICATION INFO.:

RELATED APPLN. INFO.:

Continuation of Ser. No. US 2000-602787, filed on 23 JUD 2000 DENDING

	Jun 2000, PENDING	
	NUMBER	DATE
PRIORITY INFORMATION:	DE 1999-19931454	19990708
INTONITI INTONIMITON:	DE 1999-19931478	19990708
	DE 1999-19931563	19990708
	DE 1999-19932122	19990709
	DE 1999-19932124	19990709
	DE 1999-19932125	19990709
	DE 1999-19932128	19990709
	DE 1999-19932180	19990709
	DE 1999-19932182	19990709
	DE 1999-19932190	19990709
	DE 1999-19932191	19990709
	DE 1999-19932209	19990709
	DE 1999-19932212	19990709
	DE 1999-19932227	19990709
	DE 1999-19932228	19990709
	DE 1999-19932229	19990709
	DE 1999-19932230	19990709
•	DE 1999-19932927	19990714
	DE 1999-19933005	19990714
	DE 1999-19933006	19990714
	DE 1999-19942088	19990903
	DE 1999-19940764	19990827

DE 1999-19940765 19990827 DE 1999-19940766 19990827 DE 1999-19940830 19990827 DE 1999-19940831 19990827 DE 1999-19940832 19990827 DE 1999-19940833 19990827 DE 1999-19941378 19990831 DE 1999-19941379 19990831 DE 1999-19942078 19990903 DE 1999-19941395 19990831 DE 1999-19942077 19990903 DE 1999-19942079 19990903 US 1999-141031P 19990625 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: LAHIVE & COCKFIELD, LLP., 28 STATE STREET, BOSTON, MA,

02109

NUMBER OF CLAIMS: 38
EXEMPLARY CLAIM: 1
LINE COUNT: 3058

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB Isolated nucleic acid molecules, designated MCT nucleic acid molecules, which encode novel MCT proteins from Corynebacterium glutamicum are described. The invention also provides antisense nucleic acid molecules, recombinant expression vectors containing MCT nucleic acid molecules, and host cells into which the expression vectors have been introduced. The invention still further provides isolated MCT proteins, mutated MCT proteins, fusion proteins, antigenic peptides and methods for the improvement of production of a desired compound from C. glutamicum based on genetic engineering of MCT genes in this organism.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

L5 ANSWER 111 OF 176 USPATFULL on STN

ACCESSION NUMBER: 2001:196823 USPATFULL

TITLE: SAM operon

INVENTOR(S): DeHoff, Bradley Stuart, Indianapolis, IN, United States

Rosteck, Jr., Paul Robert, Indianapolis, IN, United

States

PATENT ASSIGNEE(S): Eli Lilly and Company, Indianapolis, IN, United States

(U.S. corporation)

NUMBER DATE

PRIORITY INFORMATION: US 1996-30898P 19961113 (60)

DOCUMENT TYPE: Utility
FILE SEGMENT: GRANTED
PRIMARY EXAMINER: Yucel, Remy

LEGAL REPRESENTATIVE: Cohen, Charles E., Tucker, Tina M., Webster, Thomas D.

NUMBER OF CLAIMS: 36 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 2 Drawing Figure(s); 2 Drawing Page(s)

LINE COUNT: 725

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

The invention provides isolated nucleic acid compounds encoding a novel SAM synthetase of Streptomyces fradiae. Also provided are vectors and transformed heterologous host cells for expressing the SAM synthetase and a method for preparing S-adenosylmethionine from recombinant host cells transformed with the SAM synthetase gene.

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

ANSWER 114 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 19

ACCESSION NUMBER: 2001:64719 CAPLUS

135:148135 DOCUMENT NUMBER:

Lowering S-adenosylmethionine levels in Escherichia TITLE:

coli modulates C-to-T transition mutations

AUTHOR (S): Macintyre, Georgina; Atwood, C. Victoria; Cupples,

Claire G.

Biology Department, Concordia University, Montreal, CORPORATE SOURCE:

QC, H3G 1M8, Can.

Journal of Bacteriology (2001), 183(3), 921-927 SOURCE:

CODEN: JOBAAY; ISSN: 0021-9193

PUBLISHER: American Society for Microbiology

Journal DOCUMENT TYPE: English LANGUAGE:

Deoxycytosine methylase (Dcm) enzyme activity causes mutagenesis in vitro either directly by enzyme-induced deamination of cytosine to uracil in the absence of the Me donor, S-adenosylmethionine (SAM), or indirectly through spontaneous deamination of [5-methyl]cytosine to thymine. Using a Lac reversion assay, we investigated the contribution of the first mechanism to Dcm mutagenesis in vivo by lowering the levels of SAM.

Escherichia coli SAM levels were lowered by

reducing SAM synthetase activity via the introduction of a metK84 allele or by hydrolyzing SAM using the bacteriophage T3 SAM hydrolase. The metK84 strains exhibited increased C-to-T mutagenesis. Expression of the T3 SAM hydrolase gene, under the control of the arabinose-inducible PBAD promoter, effectively reduced Dcm-mediated genomic DNA methylation. However, increased mutagenesis was not observed until extremely high arabinose concns. were used, and genome methylation at Dcm sites was

THERE ARE 50 CITED REFERENCES AVAILABLE FOR THIS REFERENCE COUNT: 50 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 115 OF 176 BIOSIS COPYRIGHT (c) 2006 The Thomson Corporation on L5 STN

ACCESSION NUMBER: 2002:201211 BIOSIS PREV200200201211 DOCUMENT NUMBER:

metK is an essential gene in E. coli K-12. TITLE:

Wei, Y. [Reprint author]; Newman, E. [Reprint author] AUTHOR (S):

CORPORATE SOURCE: Concordia University, Montreal, PQ, Canada

Abstracts of the General Meeting of the American Society SOURCE:

for Microbiology, (2001) Vol. 101, pp. 418-419. print. Meeting Info.: 101st General Meeting of the American Society for Microbiology. Orlando, FL, USA. May 20-24,

2001. American Society for Microbiology.

ISSN: 1060-2011.

DOCUMENT TYPE: Conference; (Meeting)

Conference; Abstract; (Meeting Abstract)

LANGUAGE: English

negligible.

Entered STN: 20 Mar 2002 ENTRY DATE:

Last Updated on STN: 20 Mar 2002

The E. coli metK gene codes for the enzyme s-adenosylmethionine AB synthetase which converts methionine to S-adenosylmethionine (SAM) a major methyl donor of the cell. To see if this gene is essential to E. coli, we used the elegant pKO3 gene replacement vector of George Church in cells carrying an intact metK gene on the paraBAD plasmid. This resulted in a 600 bp deletion in metK, confirmed by sequencing across the deletion cloned in pKO3. Because this strain allowed for frequent recombination, and because pBAD-metK is expressed at a relatively high rate even in the absence of arabinose, we wished to clone the intact metK gene under the control of a highly-inducible tightly-regulated promoter in a low copy plasmid in a recA- strain. therefore cloned metK on a p15A construct obtained from Dr. Lutz, under

control of the pLtet promoter, and replaced pbadmetK with this vector. We then transduced this strain with a gene producing the tet repressor, and made it recA- also by transduction. This resulted in a strain of E. coli K12 which carries a 600 bp deletion in the chromosomal metK gene with a plasmid-carried functional metK under the tet promoter. This strain cannot grow on LB unless anhydrotetracycline is added to induce metK transcription. We conclude that metK is an essential gene, and E. coli K12 cannot grow without SAM.

ANSWER 143 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 37

ACCESSION NUMBER: 1997:108749 CAPLUS

DOCUMENT NUMBER: 126:225514

TITLE: Enzymic synthesis of S-adenosyl-L-methionine on the

preparative scale

AUTHOR (S): Park, Jeongho; Tai, Junzhe; Roessner, Charles A.;

Scott, A. Ian

Center for Biological NMR, Department of Chemistry, CORPORATE SOURCE:

Texas AandM University, College Station, TX,

77843-3255, USA

SOURCE: Bioorganic & Medicinal Chemistry (1996), 4(12),

2179-2185

CODEN: BMECEP; ISSN: 0968-0896

PUBLISHER: Elsevier DOCUMENT TYPE: Journal LANGUAGE: English

The problems inherent in the enzymic and chemical synthesis of S-adenosyl-L-methionine (SAM) led to development of an efficient, simple method for the synthesis of large amts. of labeled SAM. It has previously been reported that the problem of product inhibition of E.

coli SAM synthetase encoded by the metK gene

was successfully overcome in the presence of sodium p-toluenesulfonate (pTsONa). This research has now been expanded to demonstrate that product inhibition of this enzyme can also be overcome by adding a high concentration

of

 β -mercaptoethanol (β ME), acetonitrile, or urea. In addition, a recombinant strain of E. coli has been constructed that expresses the yeast SAM synthetase encoded by the sam2 gene. The yeast enzyme does not have the problem of product inhibition seen with the E. coli enzyme. Complete conversion of 10 mM methionine to SAM was achieved in incubations with either the recombinant yeast enzyme and 1 M potassium ion or the E. coli enzyme in the presence of additives such as βME , acetonitrile, urea, or pTsONa. The recombinant yeast SAM synthetase was used to generate SAM in situ for use in the multi-enzymic synthesis of precorrin 2.

REFERENCE COUNT:

57 THERE ARE 57 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 146 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER:

1995:605638 CAPLUS

DOCUMENT NUMBER:

123:2775

TITLE:

Production of s-adenosyl-methionine (SAM) by fermentation of Escherichia coli transformed with rat S-Adenosyl-

methionine synthetase

INVENTOR (S):

Mato, Jose Maria; Pajares, Maria Angeles; Mingorance,

Jesus; Avarez, Luis

PATENT ASSIGNEE(S):

Boehringer Ingelheim Espana S.A., Spain

SOURCE: Eur. Pat. Appl., 11 pp.

CODEN: EPXXDW

DOCUMENT TYPE:

Patent

LANGUAGE:

English

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO. KIND DATE APPLICATION NO. DATE ______

EP 647712 Al 19950412 EP 1993-116221 19931007
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LI, LU, MC, NL, PT, SE
PRIORITY APPLN. INFO.: EP 1993-116221 19931007
AB A method of production for production of S-adenosyl-L-methionine (SAM) by fermentation

of bacteria is presented. Escherichia coli was transformed with an expression vector containing a nucleic acid sequence coding for rat S-adenosyl-methionine synthetase and an inducible bacteriophage T7 promoter sequence. SAM is a naturally occurring mol., widely distributed throughout body tissues and fluids. It plays a central role in three main metabolic pathways, such as transmethylation, transsulphuration and aminopropylation. SAM is currently used in medical and pharmacol. areas due to its therapeutic potential in liver damage and affective disorders. This method has superior properties with respect to prior approaches to SAM production by fermentation of a microorganism and may have a potential industrial interest as a powerful source of SAM.

L5 ANSWER 169 OF 176 CAPLUS COPYRIGHT 2006 ACS on STN DUPLICATE 49

ACCESSION NUMBER: 1987:512408 CAPLUS

DOCUMENT NUMBER: 107:112408

TITLE: A mutant of Escherichia coli temperature sensitive in

the biosynthesis of S-adenosylmethionine

AUTHOR(S): Kimchi, Bracha; Ron, Eliora Z.

CORPORATE SOURCE: George S. Wise Fac. Life Sci., Tel Aviv Univ., Tel

Aviv, 69978, Israel

SOURCE: FEMS Microbiology Letters (1987), 43(1), 101-6

CODEN: FMLED7; ISSN: 0378-1097

DOCUMENT TYPE: Journal LANGUAGE: English

S-Adenosylmethionine (SAM) is synthesized in vitro by SAM synthetase, which is coded for by the metK gene of E. coli. Since E. coli cells are impermeable to SAM, it has been impossible to obtain mutants which require SAM for growth. The finding that SAM is required for regulating methionine biosynthesis was used to select metK mutants which have the phenotype of resistance to methionine analogs. These mutants show reduced SAM synthesis in vitro, and a lower level of intracellular SAM. However, as all these mutants can grow, without added SAM, the block in SAM synthesis must be incomplete. To try to block SAM synthesis more completely temperature-sensitive metK mutation was combined with a block in the synthesis of cystathionine (metA), a precursor that supports only slow growth because of slow entry. At the nonpermissive temperature for metK, these double mutants could grow on methionine but not on cystathionine, in correlation with the greatly reduced intracellular level of SAM. This result indicates that it is possible to reduce intracellular SAM concns. to levels which are low enough to prevent growth of E. coli. Moreover, since the mutation responsible for the reduction in SAM level is in the metK gene, it is suggested that SAM synthetase is the enzyme which catalyzes SAM biosynthesis in E. coli cells.

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INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, ANTE, AQUALINE, AQUASCI, BIOENG, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CAPLUS, CEABA-VTB, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DISSABS, DRUGB, DRUGMONOG2, DRUGU, EMBAL, EMBASE, ...' ENTERED AT 20:40:48 ON 04 SEP 2006 SEA (SAM OR ADOMET?) (S) (SYNTHETAS? OR SYNTHAS?)

⁴⁸ FILE AGRICOLA

¹ FILE AQUASCI

¹⁸ FILE BIOENG

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280
                    FILE BIOSIS
              40
                    FILE BIOTECHABS
               40
                    FILE BIOTECHDS
             158
                    FILE BIOTECHNO
                    FILE CABA
             126
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                D RANK
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     BIOTECHNO, LIFESCI, TOXCENTER' ENTERED AT 20:42:47 ON 04 SEP 2006
L2
           2227 SEA (SAM OR ADOMET?) (S) (SYNTHETAS? OR SYNTHAS?)
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